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OCT 26 2000

TECH CENTER 1600/2900



SEQUENCE LISTING

<110> Ludevid, Doloros
Torrent, Margarita
Alvarez, Inaki
Perez, Pascual

<120> Amino acid-enriched plant protein
reserves, particularly lysine-enriched maize gamma-zein, and
plants expressing such proteins

<130> 50062/004001

<140> 09/117,246
<141> 1998-12-03

<150> PCT/FR97/00167
<151> 1997-01-28

<150> FR96/01004
<151> 1996-01-29

<160> 11

<170> FastSEQ for Windows Version 4.0

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<212> DNA
<213> Artificial Sequence

<220>
<223> based on Maize

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<220>
<223> based on Maize

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<211> 17
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1 5 10 15
Pro

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1 5 10 15
Lys Pro Lys Pro Lys Pro Lys Glu Phe Leu Gln Pro
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<212> PRT
<213> Maize

<400> 5
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1 5 10 15
Phe Lys Leu Asp
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<211> 672
<212> DNA
<213> Maize

<220>
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<222> (1) ... (672)

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1 5 10 15

gcc acc tcc acg cat aca agc ggc ggc tgc ggc tgc cag cca ccg ccg 96
Ala Thr Ser Thr His Thr Ser Gly Gly Cys Gly Cys Gln Pro Pro Pro

-2
40

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Pro Val His Leu Pro Pro Pro Val His Leu Pro Pro Pro Val His Leu				
35	40	45		
cca cct ccg gtg cat ctc cca ccg ccg gtc cac ctg ccg ccg gtc				192
Pro Pro Pro Val His Leu Pro Pro Pro Val His Leu Pro Pro Pro Val				
50	55	60		
cac ctg cca ccg ccg gtc cat gtg ccg ccg gtt cat ctg ccg ccg				240
His Leu Pro Pro Pro Val His Val Pro Pro Pro Val His Leu Pro Pro				
65	70	75	80	
cca cca tgc cac tac act caa ccg ccc ccg cct cag cct cat ccc				288
Pro Pro Cys His Tyr Pro Thr Gln Pro Pro Arg Pro Gln Pro His Pro				
85	90	95		
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Gln Pro His Pro Cys Pro Cys Gln Gln Pro His Pro Ser Pro Cys Gln				
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ctg cag gga acc tgc ggc gtt ggc agc acc ccg atc ctg ggc cag tgc				384
Leu Gln Gly Thr Cys Gly Val Gly Ser Thr Pro Ile Leu Gly Gln Cys				
115	120	125		
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Val Glu Phe Leu Arg His Gln Cys Ser Pro Thr Ala Thr Pro Tyr Cys				
130	135	140		
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Ser Pro Gln Cys Gln Ser Leu Arg Gln Gln Cys Cys Gln Gln Leu Arg				
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Gln Val Glu Pro Gln His Arg Tyr Gln Ala Ile Phe Gly Leu Val Leu				
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Leu Ala Ala Gln Ile Ala Gln Gln Leu Thr Ala Met Cys Gly Leu Gln				
195	200	205		
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<213> Maize

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35 40 45
Pro Pro Pro Val His Leu Pro Pro Pro Val His Leu Pro Pro Pro Val
50 55 60
His Leu Pro Pro Pro Val His Val Pro Pro Pro Val His Leu Pro Pro
65 70 75 80
Pro Pro Cys His Tyr Pro Thr Gln Pro Pro Arg Pro Gln Pro His Pro
85 90 95
Gln Pro His Pro Cys Pro Cys Gln Gln Pro His Pro Ser Pro Cys Gln
100 105 110
Leu Gln Gly Thr Cys Gly Val Gly Ser Thr Pro Ile Leu Gly Gln Cys
115 120 125
Val Glu Phe Leu Arg His Gln Cys Ser Pro Thr Ala Thr Pro Tyr Cys
130 135 140
Ser Pro Gln Cys Gln Ser Leu Arg Gln Gln Cys Cys Gln Gln Leu Arg
145 150 155 160
Gln Val Glu Pro Gln His Arg Tyr Gln Ala Ile Phe Gly Leu Val Leu
165 170 175
Gln Ser Ile Leu Gln Gln Gln Pro Gln Ser Gly Gln Val Ala Gly Leu
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gcc acc tcc acg cat aca agc ggc ggc tgc ggc cag cca ccg ccg 96

4-
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Pro Val His Leu Pro Pro Val His Leu Pro Pro Pro Val His Leu			
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ccs cct ccg gtg cat ctc cca ccg ccg gtc cac ctg ccg ccg gtc	192		
Pro Pro Pro Val His Leu Pro Pro Pro Val His Leu Pro Pro Pro Val			
50	55	60	
cac ctg cca ccg ccg gtc cat gtg ccg ccg gtt cat ctg ccg ccg	240		
His Leu Pro Pro Pro Val His Val Pro Pro Pro Val His Leu Pro Pro			
65	70	75	80
ccs cca tgc cac tac cct act caa ccg ccc ccg atc gaa ttc aaa cca	288		
Pro Pro Cys His Tyr Pro Thr Gln Pro Pro Arg Ile Glu Phe Lys Pro			
85	90	95	
aag cca aag ccg aag cca aaa gaa ttc aaa cca aag cca aag ccg aag	336		
Lys Pro Lys Pro Lys Glu Phe Lys Pro Lys Pro Lys Pro Lys Pro Lys			
100	105	110	
ccs aaa gaa ttc ctg cag ccc ctg cag gga acc tgc ggc gtt ggc agc	384		
Pro Lys Glu Phe Leu Gln Pro Leu Gln Gly Thr Cys Gly Val Gly Ser			
115	120	125	
acc ccg atc ctg ggc cag tgc gag ttt ctg agg cat cag tgc agc	432		
Thr Pro Ile Leu Gly Gln Cys Val Glu Phe Leu Arg His Gln Cys Ser			
130	135	140	
ccg acg gcg acg ccc tac tgc tcg cct cag tgc cag tcg ttg cgg cag	480		
Pro Thr Ala Thr Pro Tyr Cys Ser Pro Gln Cys Gln Ser Leu Arg Gln			
145	150	155	160
cag tgt tgc cag cag ctc agg cag gtg gag ccg cag cac ccg tac cag	528		
Gln Cys Cys Gln Gln Leu Arg Gln Val Glu Pro Gln His Arg Tyr Gln			
165	170	175	
gcg atc ttc ggc ttg gtc ctc cag tcc atc ctg cag cag cag ccg caa	576		
Ala Ile Phe Gly Leu Val Leu Gln Ser Ile Leu Gln Gln Pro Gln			
180	185	190	
agc ggc cag gtc gcg ggg ctg ttg gcg cag ata gcg cag caa ctg	624		
Ser Gly Gln Val Ala Gly Leu Leu Ala Ala Gln Ile Ala Gln Gln Leu			
195	200	205	
acg gcg atg tgc ggc ctg cag cag ccg act cca tgc ccc tac gct gct	672		
Thr Ala Met Cys Gly Leu Gln Gln Pro Thr Pro Cys Pro Tyr Ala Ala			
210	215	220	

5-
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225 230

693

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20 25 30
Pro Val His Leu Pro Pro Pro Val His Leu Pro Pro Pro Val His Leu
35 40 45
Pro Pro Pro Val His Leu Pro Pro Pro Val His Leu Pro Pro Pro Val
50 55 60
His Leu Pro Pro Pro Val His Val Pro Pro Val His Leu Pro Pro Pro
65 70 75 80
Pro Pro Cys His Tyr Pro Thr Gln Pro Pro Arg Ile Glu Phe Lys Pro
85 90 95
Lys Pro Lys Pro Lys Pro Lys Glu Phe Lys Pro Lys Pro Lys Pro Lys
100 105 110
Pro Lys Glu Phe Leu Gln Pro Leu Gln Gly Thr Cys Gly Val Gly Ser
115 120 125
Thr Pro Ile Leu Gly Gln Cys Val Glu Phe Leu Arg His Gln Cys Ser
130 135 140
Pro Thr Ala Thr Pro Tyr Cys Ser Pro Gln Cys Gln Ser Leu Arg Gln
145 150 155 160
Gln Cys Cys Gln Gln Leu Arg Gln Val Glu Pro Gln His Arg Tyr Gln
165 170 175
Ala Ile Phe Gly Leu Val Leu Gln Ser Ile Leu Gln Gln Gln Pro Gln
180 185 190
Ser Gly Gln Val Ala Gly Leu Leu Ala Ala Gln Ile Ala Gln Gln Leu
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Ala Gly Gly Val Pro His
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<212> DNA
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 Pro Val His Leu Pro Pro Val His Leu Pro Pro Val His Leu
 35 40 45

 cca cct ccg gtg cat ctc cca ccg ccg gtc cac ctg ccg ccg ccg gtc 192
 Pro Pro Pro Val His Leu Pro Pro Val His Leu Pro Pro Pro Val
 50 55 60

 cac ctg cca ccg ccg gtc cat gtg ccg ccg gtt cat ctg ccg ccg 240
 His Leu Pro Pro Pro Val His Val Pro Pro Pro Val His Leu Pro Pro
 65 70 75 80

 cca cca tgc cac tac cct act caa ccg ccc ccg cct cag cct cat ccc 288
 Pro Pro Cys His Tyr Pro Thr Gln Pro Pro Arg Pro Gln Pro His Pro
 85 90 95

 cag cca cac cca tgc ccg tgc caa cag ccg cat cca agc ccg tgc cag 336
 Gln Pro His Pro Cys Pro Cys Gln Gln Pro His Pro Ser Pro Cys Gln
 100 105 110

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 Ile Glu Phe Lys Pro Lys Pro Lys Pro Lys Pro Lys Glu Phe Leu Gln
 115 120 125

 ccc ctg cag gga acc tgc ggc gtt ggc agc acc ccg atc ctg ggc cag 432
 Pro Leu Gln Gly Thr Cys Gly Val Gly Ser Thr Pro Ile Leu Gly Gln
 130 135 140

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 Cys Val Glu Phe Leu Arg His Gln Cys Ser Pro Thr Ala Thr Pro Tyr
 145 150 155 160

 tgc tcg cct cag tgc cag tcg ttg ccg cag cag tgt tgc cag cag ctc 528
 Cys Ser Pro Gln Cys Gln Ser Leu Arg Gln Gln Cys Cys Gln Gln Leu
 165 170 175

 agg cag gtg gag ccg cag cac ccg tac cag gcg atc ttc ggc ttg gtc 576
 Arg Gln Val Glu Pro Gln His Arg Tyr Gln Ala Ile Phe Gly Leu Val
 180 185 190

 ctc cag tcc atc ctg cag cag ccg caa agc ggc cag gtc gcg ggg 624
 Leu Gln Ser Ile Leu Gln Gln Pro Gln Ser Gly Gln Val Ala Gly

7-45

195

200

205

ctg ttg gcg gcg cag ata gcg cag caa ctg acg gcg atg tgc ggc ctg 672
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 210 215 220

cag cag ccg act cca tgc ccc tac gct gct gcc ggc ggt gtc ccc cac 720
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tga 723

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 35 40 45
 Pro Pro Pro Val His Leu Pro Pro Pro Val His Leu Pro Pro Pro Val
 50 55 60
 His Leu Pro Pro Pro Val His Val Pro Pro Pro Val His Leu Pro Pro
 65 70 75 80
 Pro Pro Cys His Tyr Pro Thr Gln Pro Pro Arg Pro Gln Pro His Pro
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 Gln Pro His Pro Cys Pro Cys Gln Gln Pro His Pro Ser Pro Cys Gln
 100 105 110
 Ile Glu Phe Lys Pro Lys Pro Lys Pro Lys Glu Phe Leu Gln
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 Pro Leu Gln Gly Thr Cys Gly Val Gly Ser Thr Pro Ile Leu Gly Gln
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 Cys Val Glu Phe Leu Arg His Gln Cys Ser Pro Thr Ala Thr Pro Tyr
 145 150 155 160
 Cys Ser Pro Gln Cys Gln Ser Leu Arg Gln Gln Cys Cys Gln Gln Leu
 165 170 175
 Arg Gln Val Glu Pro Gln His Arg Tyr Gln Ala Ile Phe Gly Leu Val
 180 185 190
 Leu Gln Ser Ile Leu Gln Gln Pro Gln Ser Gly Gln Val Ala Gly
 195 200 205
 Leu Leu Ala Ala Gln Ile Ala Gln Gln Leu Thr Ala Met Cys Gly Leu
 210 215 220
 Gln Gln Pro Thr Pro Cys Pro Tyr Ala Ala Gly Gly Val Pro His
 225 230 235 240

8
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